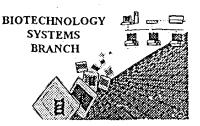
## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: $09/051/59$	
Source: 1FW16	
Date Processed by STIC: 5/ii/04 ~	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.1 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual cPAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to (EFFECTIVE 12/01/03):
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two. 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 4B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03



IFW16

RAW SEQUENCE LISTING

DATE: 05/11/2004

PATENT APPLICATION: US/09/051,159

TIME: 15:32:56

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\05112004\I051159.raw

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4 <110> APPLICANT: Balmain, Allan
                  Zhu, Jingde
        7 <120> TITLE OF INVENTION: Antitumour vector constructs and methods
      10 <130> FILE REFERENCE: CCI-005USRCE
      12 <140> CURRENT APPLICATION NUMBER: 09/051,159
      13 <141> CURRENT FILING DATE: 1999-01-13
      15 <150> PRIOR APPLICATION NUMBER: GB 9520079.6
      16 <151> PRIOR FILING DATE: 1995-10-02
      18 <150> PRIOR APPLICATION NUMBER: GB 9616685.5
      19 <151> PRIOR FILING DATE: 1996-08-08
      21 <150> PRIOR APPLICATION NUMBER: PCT/GB96/02416
      22 <151> PRIOR FILING DATE: 1996-10-02
                                                                                          Does Not Comply
      24 <160> NUMBER OF SEQ ID NOS: 3
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      26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
      28 <210> SEQ ID NO: 1
      29 <211> LENGTH: 11
      30 <212> TYPE: DNA
      31 <213> ORGANISM: Artificial Sequence
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38 <223> OTHER INFORMATION: N may be G or A

40 <221> misc_feature
41 <222> LOCATION: 5, 6, 7

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45 <222> LOCATION: (7)...(9) (9)...(11) ( no n's purent. Using w is sufficient.
46 <223> OTHER INFORMATION: N may be T or C no n's purent. Using y is sufficient.
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48 <400> 1

49 riccwwwgyy y
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      63 <211> LENGTH: 20
      64 <212> TYPE: DNA
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/051,159

DATE: 05/11/2004

TIME: 15:32:56

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\05112004\I051159.raw

67 <220> FEATURE:

68 <223> OTHER INFORMATION: Synthetic construct

70 <400> SEQUENCE: 3

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## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/051,159

DATE: 05/11/2004 TIME: 15:32:57

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\05112004\I051159.raw

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L:40 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:44 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:48 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1